

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/535, 523 A  
Source: PCT  
Date Processed by STIC: 09/26/2005

# ***ENTERED***



PCT

## RAW SEQUENCE LISTING

DATE: 09/26/2005

PATENT APPLICATION: US/10/535,523A

TIME: 12:50:39

Input Set : D:\seq listing.app

Output Set: N:\CRF4\09262005\J535523A.raw

3 &lt;110&gt; APPLICANT: The Government of the United States of America

5 &lt;120&gt; TITLE OF INVENTION: CELL LINES AND HOST NUCLEIC ACIDS RELATED TO INFECTIOUS

## DISEASES

7 &lt;130&gt; FILE REFERENCE: 6395-66741-06

C--&gt; 9 &lt;140&gt; CURRENT APPLICATION NUMBER: US/10/535,523A

C--&gt; 9 &lt;141&gt; CURRENT FILING DATE: 2005-05-17

9 &lt;150&gt; PRIOR APPLICATION NUMBER: PCT/03/037143

10 &lt;151&gt; PRIOR FILING DATE: 2003-11-18

12 &lt;150&gt; PRIOR APPLICATION NUMBER: US 60/482,604

13 &lt;151&gt; PRIOR FILING DATE: 2003-06-25

15 &lt;150&gt; PRIOR APPLICATION NUMBER: US 60/427,464

16 &lt;151&gt; PRIOR FILING DATE: 2002-11-18

18 &lt;160&gt; NUMBER OF SEQ ID NOS: 845

20 &lt;170&gt; SOFTWARE: PatentIn version 3.2

22 &lt;210&gt; SEQ ID NO: 1

23 &lt;211&gt; LENGTH: 937

24 &lt;212&gt; TYPE: DNA

25 &lt;213&gt; ORGANISM: Homo sapiens

28 &lt;220&gt; FEATURE:

29 &lt;221&gt; NAME/KEY: misc\_feature

30 &lt;222&gt; LOCATION: (1)..(937)

31 &lt;223&gt; OTHER INFORMATION: n is a, g, c, or t

33 &lt;400&gt; SEQUENCE: 1

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 36 cgaaacgata ctyrgstacc gcttgccaaa cctacagggtg ggggtctttca agagggtctcc 120  
 38 agacctaggg gagcatctca gcgtcactcg ctgtccagtt gctgtgatca ggtgctttgg 180  
 40 ggtttgtgtg actccagaat ccactgggccc tgtgtgtcag aagacaaaag ttaaccataa 240  
 42 ggcacagaag aaagcctcct gctgaagcca tcggtggccc acatgcattt cagggacaag 300  
 44 aaatgaagat cggagacttt caagttgtgc ccaggactca cctgctccca ggagacaaaa 360  
 46 ggccacacag cagaggagcc tgaagcccat ggcaggatct cctagcttgg ggctggtgtc 420  
 48 tctgtagtaa gcattctgaa gttcctaagc tcccttcttc ctgataggag cattgacctg 480  
 50 tgatgtcacc aactgacat actttccctt gcaggccact ccagccact gtactctttg 540  
 52 gcaggcctca ggttctgcta ctccatgtac tattctgtc ttgcacaggc cagaagctaa 600  
 54 aggtgaggag gactgaacac agtaccaaca taccacatc acaccttact ttcctctgcc 660  
 56 cgccctgtcc ctgccctgac actgattccc cagcccttgc caccacagcc ccttcaccct 720  
 58 ccactgcccg tgcagcagca gagacactcc ctcccttgatg caaactgagg cctctggcac 780  
 60 cccaactctt tcaaggcaat gatagtctgt gcttaactct acatggccag gccccactc 840  
 62 aggggaattnc tgtgtgaaat tgttatccgc tgsacaattc cacacaacat ggnncgtcag 900  
 64 accccgaaga aaagaancaa nggatctttt ggnnacc 937

67 &lt;210&gt; SEQ ID NO: 2

68 &lt;211&gt; LENGTH: 1515

69 &lt;212&gt; TYPE: DNA

70 &lt;213&gt; ORGANISM: Homo sapiens

73 &lt;220&gt; FEATURE:

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76 <223> OTHER INFORMATION: n is a, c, g, or t
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83 tatgcagtgc tgccataacc atgagtata acactgcggc caacttactt ctgacaacga      180
85 tcggaggacc gaaggagcta accgcttttt tgcacaacat gggggatcat gtaactcgcc      240
87 ttgatcgttg ggaaccggag ctgaatgaag ccataccaaa cgacgagcgt gacaccacga      300
89 tgcctgcagc aatggcaaca acgttgcgca aactattaac tggcgaacta cttactctag      360
91 cttcccggca acaattaata gactggatgg aggcggataa agttgcagga ccacttctgc      420
93 gtcgggccct tccggctggc tggtttattg ctgataaatc tggagccggg gagcgtgggt      480
95 ctgcgggtat cattgcagca ctggggccag atggtaagcc ctcccgtatc gtagttatct      540
97 acacgacggg gagtcaggca actatggatg aacgaaatag acagatcgct gagatagggtg      600
99 cctcactgat taagcattnng gkaanctgtc agaccattkt ttactgcata tacsgatcca      660
101 ttgcccttat ctcaaactct tattatgaaa tcactnccct tgagagaraa aaagcctttt      720
103 tctcttnngg atkgtcccag magytyccga mcatcccac tycccaacct tatgkggccc      780
105 agcaatgans cctagtagta ggaaaatcty tatggatacy ggkgnctgak ggggaarattc      840
107 ttcytctcat gaarwgatgg kgactggggc tytgggatgc tcacgggaat cctattttcc      900
109 cccacaaaga agttatttta ttacacaacc atttgatga cccctttttt cttccaattn      960
111 nccaaataaa tctgtaaagg tcacaggtga agttcttctc ttttaagagct actccatgct      1020
113 aagttcagcg agaacttggg gtaccctaga cattcttcca gagatgcttt tcttgtaact      1080
115 cttttcaata agtaagcatg ctttgcctctg cactgggtgt cactgtgtgt ggatgctgtt      1140
117 gtccctgcct tgccctatat tctgtccaca tggtttcttc ataggatgat gcttaggtca      1200
119 gccctgaggt ttgaaccagt caacaagtcc aggttgggtg ggagtccctt tagtacctcc      1260
121 ctttgcagga ataatgctgc acccagaaac tccctcagag cctctccact gggagggggcc      1320
123 ttgtgaccat tcctggttta ctctcttctg tccagcatcc catgtggcca atggggccct      1380
125 ttcattttca atggatatctc anttnttaca gtaagttata ttattgccct acatngaact      1440
127 catctttttt cantgttacc tgnngaagaa tggnaagga tgcccnnaan tnggcccaaa      1500
129 anaatccact tcgnn                                     1515
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133 <211> LENGTH: 885
134 <212> TYPE: DNA
135 <213> ORGANISM: Homo sapiens
138 <220> FEATURE:
139 <221> NAME/KEY: misc_feature
140 <222> LOCATION: (1)..(885)
141 <223> OTHER INFORMATION: n is a, g, c, or t
143 <400> SEQUENCE: 3
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146 cgagttcttc tgagcgggac tctgggggtc gaaatgagct agcccttaag taacgccatt      120
148 ttgcaaggca tggaaaaata cataactgag aatagaaaag ttcagatcga ggtcaggaac      180
150 agatggaaca gggtcgaccg gtcgaccggt cgaccctaga gaaccatcag atgtttccag      240
152 ggtgccccaa ggacctgaaa tgacctgtg ccttatttga actaaccaat cagtctcgtt      300
154 ctgcgttctg ttgcgcgct tctgtctccc gagctcaata aaagagccca caaccctca      360
156 ctcgggggcg cagtctccg attgactgag tcgcccgggt acccgtgtat ccaataaacc      420
158 ctcttgcaat tgcacccgac ttgtggtctc gctgttccct gggaggggtct cctctgagt      480
160 attgactacc cgtcagcggg ggtctttcac tctctgtgta ctggtaccaa cagagcctgg      540
162 accagggcct ccagttctc attcagtatt ataattggaga agagagagca aaaggaaaca      600

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164 ttcttgaacg attctccgca caacagttcc ctgacttgca ctctgaacta aacctgagct      660
166 ctctggagct gggggactca gctttgtatt tctgtgccag cagcgtaggt ggtagcttga      720
168 aacagttctt cggggcaggg acacggctca ccgtgctagg taagaagggg gctccagtgg      780
170 gagagagggg gagcagccca nccctgnncca ccccanancc tgttnttagg ggagtggnc      840
172 ctgggcatcc aggcctnct cnaggaancc gggtncgcn ggncc                        885
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176 <211> LENGTH: 900
177 <212> TYPE: DNA
178 <213> ORGANISM: Homo sapiens
181 <220> FEATURE:
182 <221> NAME/KEY: misc_feature
183 <222> LOCATION: (1)..(900)
184 <223> OTHER INFORMATION: n is a, g, c, or t
186 <400> SEQUENCE: 4
187 ttggttaactg tcagaccaag ttactcata tcggatccag ttggagccat aagtcgtcag      60
189 acatagaaaa aaatctgaaa agatatctca aaagcccaga catttattca cactaacggg      120
191 gaaaagcata cccacagtg tcagtggagg caacatgggg tcctggattt cctcttcacc      180
193 ctcagtggta gtgaggtggt cctctcactc cttctgagta gaggaagcca agaggaaagc      240
195 tggaacttgt accatcatcc agtggtgata aagcctctgt ccctccacct tacccccagg      300
197 ttatcagtgg caaccacatg gctagtggta cccctccgcg tcctagccag aatgatata      360
199 gcagaggcct agagagtagc ccaaaaactc atctgcacc agcaggactg aggtttccta      420
201 cccccaccaa tggaagccaa gtgaggaacc taagccttca cctctcactc agcaggaacc      480
203 agacaacacc ccctaacaca cacacacaca cacacacaca cacacccttc tgttagtg      540
205 gtatcaagga ggcttgataa aatagaagat ttaaatagga tccattgccc ttatctcaa      600
207 ctcttattat gaaatcactc ccttgagaga gaaaaagcc ttttctctt ggattgtccc      660
209 agcagctccc gaccatcccc actcccaac cttatgtggc ccagcaatg agcctagtag      720
W--> 211 taggaaaatc tctatggata ctggtgctga tgggaagatt ctctctctca ngaagtgatg      780
213 gtgactgggg ctctgggatg ctcacgggaa tnccatttcc cccacaagaa nttattttat      840
215 naccaaccat ttgatgacc ccttttntt ccatttncca annaatttgt aaggncaag      900
218 <210> SEQ ID NO: 5
219 <211> LENGTH: 869
220 <212> TYPE: DNA
221 <213> ORGANISM: Homo sapiens
224 <220> FEATURE:
225 <221> NAME/KEY: misc_feature
226 <222> LOCATION: (1)..(869)
227 <223> OTHER INFORMATION: n is a, g, c, or t
229 <400> SEQUENCE: 5
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232 ttcttccaat ttccaaata aatctgtaa ggtcacagg gaagttcttc tctttaagag      120
234 ctactccatg ctaagttcag cgagaacttg ggtacccta gacattcttc cagagatgct      180
236 tttcttgtaa ctcttttcaa taagtaagca tgctttgctc tgcactgggt gtcacctgtg      240
238 ttgatgctg ttgtccctgc cttgccctat attctgtcca catggtttct tcataggatg      300
240 atgcttaggt cagccctgag gtttgaacca gtcaacaagt ccaggttggg gtggagtccc      360
242 tttagtacct ccctttgcag gaataatgct gcaccagaa actccctcag agcctctcca      420
244 ctggaggggc cttgtgacca ttctgggtt actcctcttg ttccagcatc ccatgtggcc      480
246 aatgggcccc ttctattttc aatggtatct caattcttac agtaagttat attattgccc      540
248 tacatcgaac tcatttttct tcagtgttac ctgaggaaga atggagagga tgcccagaat      600
250 tggcccagaa gaatccactt cgattctaga gaaaaaggca ggtagaggca gaagagattc      660

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Input Set : D:\seq\_listing.app

Output Set: N:\CRF4\09262005\J535523A.raw

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254 ctgtaaaaact tgggaagagg aacctgccct ggggtcaagta ggggtgttggg aggaccagat 780
256 ggagcttgaa gctctctcca tctttgtcaa gtcccctgga ctgagagggg aaaatnacat 840
258 ggccctttatc ctccagagga aantnatc 869
261 <210> SEQ ID NO: 6
262 <211> LENGTH: 850
263 <212> TYPE: DNA
264 <213> ORGANISM: Homo sapiens
267 <220> FEATURE:
268 <221> NAME/KEY: misc_feature
269 <222> LOCATION: (1)..(850)
270 <223> OTHER INFORMATION: n is a, g, c, or t
272 <400> SEQUENCE: 6
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275 gagttcttct gagggggact ctgggggttcg aaatgagcta gcccttaagt aacgccattt 120
277 tgcaaggcat ggaaaaatac ataactgaga atagaaaagt tcagatcgag gtcaggaaca 180
279 gatggaacag ggtcgaccgg tcgaccgggc gacctagag aaccatcaga tgtttccagg 240
281 gtgccccaa gacctgaaat gacctgtgc cttatttgaa ctaaccaatc agttcgcttc 300
283 tcgcttctgt tcgcgcgctt ctgctccccg agctcaataa aagagccac aaccctcac 360
285 tcggggcgcc agtcctccga ttgactgagt cgcccggtta cccgtgtatc caataaacc 420
287 tcttgagtt gcatccgact tgtggtctcg ctgttccttg ggagggtctc ctctgagtga 480
289 ttgactaccc gtcagcgggg gtctttcact ctctgtgtac tggtaacca agagcctgga 540
291 ccagggcctc cagttcctca ttcagtatta taatggagaa gagagagcaa aaggaaacat 600
293 tcttgaacga ttctccgcac aacagttccc tgacttgac tctgaactaa acctgagctc 660
295 tctggagctg ggggactcag ctttgtattt ctgtgccagc agcgtaggtg gtagcttgaa 720
297 acagttcttc gggccaggga cacggctcac cgtgctaggt aagaaggggg ctccaggtgg 780
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301 actgggncat 850
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305 <211> LENGTH: 847
306 <212> TYPE: DNA
307 <213> ORGANISM: Homo sapiens
310 <220> FEATURE:
311 <221> NAME/KEY: misc_feature
312 <222> LOCATION: (1)..(847)
313 <223> OTHER INFORMATION: n is a, g, c, or t
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320 actccagaat cactggggc tgtgtgtcag aagacaaaag ttaaccataa ggcacagaag 180
322 aaagcctcct gctgaagcca tcgttggccc acatgcattt cagggacaag aaatgaagat 240
324 cggagacttt caagttgtgc ccaggactca cctgctccca ggagacaaaa ggccacacag 300
326 cagaggagcc tgaagcccat ggcaggatct cctagcttgg ggctgggtgc tctgtagtaa 360
328 gcattctgaa gttccctaagc tcccttcttc ctgataggag cattgacctg tgatgtcacc 420
330 aactgacat actttccctt gcaggccact ccagccact gtactctttg gcaggcctca 480
332 ggttctgcta ctccatgtac tattctgtc ttgcacaggg cagaagctaa aggtgaggag 540
334 gactgaacac agtaccaaca taccacatc acaccttact ttctctgcc cgccctgtcc 600
336 ctgccctgac actgattccc cagcccttgc caccacagcc ccttcacct ccactgcccg 660
338 tgcagcagca gagacactcc ctcttgatg caaactgagg cctctggcac cccaactctt 720

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Output Set: N:\CRF4\09262005\J535523A.raw

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340 tcagggcaat gatagtctgt gcttaactct acatggccag gccccactca ggggaattctc 780
W--> 342 acctagaatt tcatatncag ccaaactaag cttcataagt gaaggggaaa taaatgctt 840
344 tacagac 847
347 <210> SEQ ID NO: 8
348 <211> LENGTH: 755
349 <212> TYPE: DNA
350 <213> ORGANISM: Homo sapiens
353 <220> FEATURE:
354 <221> NAME/KEY: misc_feature
355 <222> LOCATION: (1)..(755)
356 <223> OTHER INFORMATION: n is a, g, c, or t
358 <400> SEQUENCE: 8
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363 caggctagag tgcagtggca tgatcttggc tcaactgcaac ctccacctcc caggttcaag 180
365 tgattctctt gcctcagcct cccaagtagc tagtattaca gacgcctgcc accacgcccg 240
367 gttaattttt gtacttttag tagagacagg tttcaccata ttggccaggc tggctctcaa 300
369 ctctgacact caggtgatcc tctgcctca gcctcccaaa gtgctgggat tacaggcatg 360
371 agctaccacg tctggcctaa gtgcattgta cctataactaa caaaaccaca cttctgcctc 420
373 gaatgagaac agtctcctga acatcttgcc tctttgcctg actcaaagcc tcagggtctaa 480
375 gcctcccat aatttctagt ctccagcaga agatcaatga caggagactc tccagggtgat 540
377 gaaattaacc aattaagtaa cctgggttgg catcctcccg tttgttcacc agctcacctn 600
379 ctgccacagg tatatccttt ctctcancca tatatgcaca aacccccctnc ccacggnaca 660
381 catannaana atttgaaga ctanaaaatc aggcanggtt tancncacct tngnggctgg 720
383 agtatggnan cctgggcccgg nacatncata cattg 755
386 <210> SEQ ID NO: 9
387 <211> LENGTH: 839
388 <212> TYPE: DNA
389 <213> ORGANISM: Homo sapiens
392 <220> FEATURE:
393 <221> NAME/KEY: misc_feature
394 <222> LOCATION: (1)..(839)
395 <223> OTHER INFORMATION: n is a, g, c, or t
397 <400> SEQUENCE: 9
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400 accttacta caaggatatt acaacaggac attttttaaa acctcaaaca tcacaaaaat 120
402 ttctaagtgc aagtttattt ttattttttt tttttttttt gagacagagt ctgctctgt 180
404 caccagagct agagtgcagt ggcatgatct tggctcactg caacctccac ctcccagggt 240
406 caagtgattc tcttgctca gcctcccaag tagctagtag tacagacgcc tgccaccacg 300
408 cccggttaat ttttgactt ttagtagaga caggtttcac catattggcc aggctggtct 360
410 caaactcctg acctcagggt atcctcctgc ctccagcctcc caaagtgctg ggattacagg 420
412 catgagctac cacgtctggc ctaagtgcac gttacctata ctaacaaaac cacacttctg 480
414 cctcgaatga gaacagtctc ctgaacatct tgccctcttg cctgactcaa agcctcagg 540
416 ctaagcctcc ccataatttc tagtctcagc agaaagatca atgacaggag actctccagg 600
418 tgatgaaatt aaccaattaa gtaacctggg ttggcatcct cccgtttgtt caccagctca 660
420 cctcctgcca caggtatata ctttctctca gccatatatg cacaaacccc ctccccacgg 720
422 cacacataga aanaatttgg aagactagaa aatcaggcna gggnttanca caccttngag 780
424 ggctggagta tggnanccng ggnccgggan atncatncnn tngaaaactt gactatggg 839
427 <210> SEQ ID NO: 10

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/535,523A

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Input Set : D:\seq listing.app

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:2; N Pos. 22,619,625,696,727,728,789,825,960,961,1402,1405,1435,1453  
Seq#:2; N Pos. 1463,1464,1474,1475,1486,1490,1492,1502,1514,1515  
Seq#:3; N Pos. 9,803,806,807,816,818,825,838,858,862,868,875,880,883  
Seq#:4; N Pos. 771,812,831,841,863,877,882,883,895  
Seq#:5; N Pos. 6,8,21,820,836,863,865  
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Seq#:7; N Pos. 797  
Seq#:8; N Pos. 7,15,600,627,649,657,665,666,669,684,696,700,703,705,713,728  
Seq#:8; N Pos. 730,741,746  
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Seq#:14; N Pos. 11,27,40,700,706,716,745,749,761,790  
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Seq#:19; N Pos. 709,712,715,721,733,739,745,747,758,759,763,764,766,768  
Seq#:20; N Pos. 6,8,733,746,752,759,760,790,803,807,821,822,823,828,830,847  
Seq#:20; N Pos. 848,856,860,861,869,872,874,876,881,882,895,912  
Seq#:21; N Pos. 2,7,36,723,724,778,799,817,1461,1553,1586,1597,1599,1604  
Seq#:22; N Pos. 8,22,712,798,837  
Seq#:23; N Pos. 8,13,14,22,44,47,54,73,76,87,117,129,1519,1534,1539,1547  
Seq#:23; N Pos. 1548,1550,1558,1560  
Seq#:24; N Pos. 498,503,506,652,655,656,658,794,821,834,858,1335,1368,1375  
Seq#:24; N Pos. 1402,1435,1437  
Seq#:25; N Pos. 8,9,15,32,680,683,700,732,736,745,754,763,764,774,787,797  
Seq#:25; N Pos. 800,805,809,814,815,816,819,830  
Seq#:26; N Pos. 5,740,744,746,770,778,784,785,795,802,820,827,839,842,845  
Seq#:26; N Pos. 852,854  
Seq#:27; N Pos. 6,7,234,235,236,797,807,822,830,839,845,864,870,873  
Seq#:28; N Pos. 3,9,13,16,17,18,19,20,21,24,27,30,31,32,33,34,39,43,47,49  
Seq#:28; N Pos. 50,51,54,56,57,798,806,820,833,853,855,864,865,878,880,884

RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 09/26/2005  
PATENT APPLICATION: US/10/535,523A      TIME: 12:50:40

Input Set : D:\seq listing.app  
Output Set: N:\CRF4\09262005\J535523A.raw

Seq#:28; N Pos. 885,891,894,900,901  
Seq#:29; N Pos. 2,8,9,18,23,26,35,41,50,809,813,826  
Seq#:30; N Pos. 2,3,4,14,18,19,20,27,28,29,31,32,34,38,41,48,49,50,65,72,85  
Seq#:30; N Pos. 90,99,771,782,789,822,827,848,858,862,872,873,879,881,884  
Seq#:30; N Pos. 885,888,889  
Seq#:31; N Pos. 12,21,574,604,688,691,713,715  
Seq#:32; N Pos. 6,250

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:238,239,240,241



## VERIFICATION SUMMARY

DATE: 09/26/2005

PATENT APPLICATION: US/10/535,523A

TIME: 12:50:40

Input Set : D:\seq listing.app

Output Set: N:\CRF4\09262005\J535523A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No  
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:34 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
M:341 Repeated in SeqNo=1  
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
M:341 Repeated in SeqNo=2  
L:144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
M:341 Repeated in SeqNo=3  
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:720  
M:341 Repeated in SeqNo=4  
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
M:341 Repeated in SeqNo=5  
L:299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:780  
M:341 Repeated in SeqNo=6  
L:342 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:780  
L:359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
M:341 Repeated in SeqNo=8  
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
M:341 Repeated in SeqNo=9  
L:439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
M:341 Repeated in SeqNo=10  
L:480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
M:341 Repeated in SeqNo=11  
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
M:341 Repeated in SeqNo=12  
L:556 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0  
M:341 Repeated in SeqNo=13  
L:595 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0  
M:341 Repeated in SeqNo=14  
L:636 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
M:341 Repeated in SeqNo=15  
L:677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0  
M:341 Repeated in SeqNo=16  
L:716 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0  
M:341 Repeated in SeqNo=17  
L:757 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0  
M:341 Repeated in SeqNo=18  
L:796 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0  
M:341 Repeated in SeqNo=19  
L:835 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0  
M:341 Repeated in SeqNo=20  
L:880 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0  
M:341 Repeated in SeqNo=21  
L:947 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
M:341 Repeated in SeqNo=22  
L:990 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
M:341 Repeated in SeqNo=23  
L:1073 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:480

**VERIFICATION SUMMARY**

DATE: 09/26/2005

PATENT APPLICATION: US/10/535,523A

TIME: 12:50:40

Input Set : D:\seq listing.app

Output Set: N:\CRF4\09262005\J535523A.raw

M:341 Repeated in SeqNo=24

L:1120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0

M:341 Repeated in SeqNo=25

L:1161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0